

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/744,882

DATE: 12/04/98
 TIME: 18:19:51

INPUT SET: S30182.raw

**This Raw Listing contains the General
 Information Section and up to the first 5 pages.**

#14

SEQUENCE LISTING

(1) General Information:

(i) APPLICANT: ISNER, Jeffrey
 ASAHARA, Takayuki

(ii) TITLE OF INVENTION: METHODS FOR REGULATING ANGIOGENESIS
 (as amended)

(iii) NUMBER OF SEQUENCES: 2

(iv) CORRESPONDENCE ADDRESS:

(A) ADDRESSEE: DIKE, BRONSTEIN, ROBERTS & CUSMAN, LLP
 (B) STREET: 130 Water Street
 (C) CITY: Boston
 (D) STATE: MA
 (E) COUNTRY: USA
 (F) ZIP: 02109

(v) COMPUTER READABLE FORM:

(A) MEDIUM TYPE: Diskette
 (B) COMPUTER: IBM Compatible
 (C) OPERATING SYSTEM: DOS
 (D) SOFTWARE: FastSEQ for Windows Version 2.0

(vi) CURRENT APPLICATION DATA:

(A) APPLICATION NUMBER: US/08/744,882
 (B) FILING DATE: 08-NOV-1996
 (C) CLASSIFICATION: 424

(vii) PRIOR APPLICATION DATA:

(A) APPLICATION NUMBER:
 (B) FILING DATE:

(viii) ATTORNEY/AGENT INFORMATION:

(A) NAME: Resnick, David S
 (B) REGISTRATION NUMBER: 34,235
 (C) REFERENCE/DOCKET NUMBER: 46963

(ix) TELECOMMUNICATION INFORMATION:

(A) TELEPHONE: 617-523-3400
 (B) TELEFAX: 617-523-6440
 (C) TELEX:

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PATENT APPLICATION US/08/744,882DATE: 12/04/98
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47

48

49

(2) INFORMATION FOR SEQ ID NO:1:

50

51

(i) SEQUENCE CHARACTERISTICS:

52

(A) LENGTH: 30 base pairs

53

(B) TYPE: nucleic acid

54

(C) STRANDEDNESS: single

55

(D) TOPOLOGY: linear

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58

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

59

60

AAGACATTTT CGGGCTCACG CTGCGCACCC

30

61

62

(2) INFORMATION FOR SEQ ID NO:2:

63

64

(i) SEQUENCE CHARACTERISTICS:

65

(A) LENGTH: 30 base pairs

66

(B) TYPE: nucleic acid

67

(C) STRANDEDNESS: single

68

(D) TOPOLOGY: linear

69

70

71

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

72

73

TGGGGTAGGC ACTTTAGTAG TTCTCCTAAC

30

74

75

PAGE: 1

SEQUENCE VERIFICATION REPORT
PATENT APPLICATION US/08/744,882

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Line	Error	Original Text
30	Wrong Classification	(C) CLASSIFICATION:424